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## SEQUENCE LISTING

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<110> GUERIN-MARCHAND, CLAUDINE
     DRUILHE, PIERRE
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<141> 2001-07-10
<150> 08/098,327
<151> 1993-11-24
<150> PCT/FR92/00104
<151> 1992-02-05
<150> FR 91 01286
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Glu Leu Pro Ser Glu As<br/>n Glu Arg Gly Tyr Tyr Ile Pro His Gl<br/>n Ser 35 40 45

Ser Leu Pro Gln Asp Asn Arg Gly Asn Ser Arg Asp Ser Lys Glu Ile 50 55 60

Ser Ile Ile Glu Lys Thr Asn Arg Glu Ser Ile Thr Thr Asn Val Glu 65 70 75 80

Gly Arg Arg Asp Ile His Lys Gly His Leu Glu Glu Lys Lys Asp Gly 85 90 95

Ser Ile Lys Pro Glu Gln Lys Glu Asp Lys Ser

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Leu Gln Glu Gln Arg Asp Leu Glu Gln Arg Lys Ala Asp Thr Lys

1 10 15

Lys Asn Leu Glu Arg Lys Lys Glu His Gly Asp Ile Leu Ala Glu Asp 20 25 30

Leu Tyr Gly Arg Leu Glu Ile Pro Ala Ile Glu Leu Pro Ser Glu Asn 35 40 45

Glu Arg Gly Tyr Tyr Ile Pro His Gln Ser Ser Leu Pro Gln Asp Asn 50 55 60

Arg Gly Asn Ser Arg Asp Ser Lys Glu Ile Ser Ile Ile Glu Lys Thr 65 70 75 80

Asn Arg Glu Ser Ile Thr Thr Asn Val Glu Gly Arg Arg Asp Ile His
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Lys Gly His Leu Glu Glu Lys Lys Asp Gly Ser Ile Lys Pro Glu Gln 100 105 110

Lys Glu Asp Lys Ser 115

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Ala Glu Asp Leu Tyr Gly Arg Leu Glu Ile Pro
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Gln Arg Lys Ala Asp Thr Lys Lys
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Glu Ser Ile Thr Thr Asn Val Glu Gly Arg Arg Asp Ile His Lys
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                                 25
Phe Asn Ser Leu Val Lys Ser Val Gln Gln Gln Gln His Asn Val
                            40
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Glu Glu Lys Val Glu Glu Ser Val Glu Glu Asn Asp Glu Glu Ser Val

Glu Glu Asn Val Glu Glu Asn Val Glu Glu Asn Asp Asp Gly Ser Val
65 70 75 80

Ala Ser Ser Val Glu Glu Ser Ile Ala Ser Ser Val Asp Glu Ser Ile 85 90 95

Asp Ser Ser Ile Glu Glu Asn Val Ala Pro Thr Val Glu Glu Ile Val

Ala Pro Thr Val Glu Glu Ile Val Ala Pro Ser Val Val Glu Lys Cys 115 . 120 125

Ala Pro Ser Val Glu Glu Ser Val Ala Pro Ser Val Glu Glu Ser Val 130 135 140

Ala Glu Met Leu Lys Glu Arg 145 150

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Phe Asn Ser Leu Val Lys Ser Val Gln Gln Gln Gln His Asn 35 40 45

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Val Lys Glu Asn Ile Leu Glu Glu Ser Gln 20 25

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1 10 15

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Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Arg Ala Lys Glu Lys Leu

- Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Arg Ala Lys Glu Lys
  65 70 75 80
- Leu Gln Glu Gln Ser Asp Leu Glu Gln Asp Arg Leu Ala Lys Glu
  85 90 95
- Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Arg Ala Lys 100 105 110
- Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Arg Ala 115 120 125
- Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Leu 130 135 140
- Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg 145 150 155 160
- Arg Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu 165 170 175
- Arg Arg Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln 180 185 190
- Glu Arg Arg Ala Lys Glu Lys Leu Gln Glu Gln Gln Arg Asp Leu Glu 195 200 205
- Gln Arg Lys Ala Asp Thr Lys Lys Asn Leu Glu Arg Lys Lys Glu His 210 215 220
- Gly Asp Ile Leu Ala Glu Asp Leu Tyr Gly Arg Leu Glu Ile Pro Ala 225 230 235 240
- Ile Glu Leu Pro Ser Glu Asn Glu Arg Gly Tyr Tyr Ile Pro His Gln
  245 250 255
- Ser Ser Leu Pro Gln Asp Asn Arg Gly Asn Ser Arg Asp Ser Lys Glu 260 265 270
- Ile Ser Ile Ile Glu Lys Thr Asn Arg Glu Ser Ile Thr Thr Asn Val 275 280 285
- Glu Gly Arg Arg Asp Ile His Lys Gly His Leu Glu Glu Lys Lys Asp 290 295 300
- Gly Ser Ile Lys Pro Glu Gln Lys Glu Asp Lys Ser 305 310 315

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agttgcaaga acaacaaagc gatttagaac aagatagact tgctaaagaa aagttacaag 300
agcagcaaag cgatttagaa caagagagac gtgctaaaga aaagttgcaa gaacaacaaa 360
gcgatttaga acaagagaga cgtgctaaag aaaagttgca agaacaacaa agcgatttag 420
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qacqtgctaa agaaaagttg caagaacaac aaagcgattt agaacaagag agacgtgcta 540
aagaaaagtt gcaagaacaa caaagcgatt tagaacaaga gagacgtgct aaagaaaagt 600
tgcaagagca gcaaagagat ttagaacaaa ggaaggctga tacgaaaaaa aatttagaaa 660
gaaaaaagga acatggagat atattagcag aggatttata tggtcgttta gaaataccag 720
ctatagaact tccatcagaa aatgaacgtg gatattatat accacatcaa tcttctttac 780
ctcaggacaa cagagggaat agtagagatt ccaaggaaat atctataata gaaaaaacaa 840
atagagaatc tattacaaca aatgttgaag gacgaaggga tatacataaa ggacatcttg 900
aagaaaagaa agatggttca ataaaaccag aacaaaaaga agataaatct
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aqtqttcaac aagaacaaca acacaatgtt gaagaaaaag ttgaagaaag tgtagaagaa 180
aatgacgaag aaagtgtaga agaaaatgta gaagaaaatg tagaagaaaa tgacgacgga 240
agtgtagcct caagtgttga agaaagtata gcttcaagtg ttgatgaaag tatagattca 300
agtattgaag aaaatgtagc tccaactgtt gaagaaatcg tagctccaac tgttgaagaa 360
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aaaaagatga aatcataaaa tctaacttga gaagtggttc ttcaaattct aggaatcgaa 180
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gaa atc ata aaa tct aac ttg aga agt ggt tct tca aat tct agg aat Glu Ile Ile Lys Ser Asn Leu Arg Ser Gly Ser Ser Asn Ser Arg Asn 35 40 45	144												
cga ata aat gag gaa aat cac gag aag aaa cac gtt tta tct cat aat Arg Ile Asn Glu Glu Asn His Glu Lys Lys His Val Leu Ser His Asn 50 55 60	192												
tca tat gag aaa act aaa aat aat gaa aat aat aaa ttt ttc gat aag Ser Tyr Glu Lys Thr Lys Asn Asn Glu Asn Asn Lys Phe Phe Asp Lys 65 70 75 80	240												
gat aaa gag tta acg atg tct aat gta aaa aat gtg tca caa aca aat Asp Lys Glu Leu Thr Met Ser Asn Val Lys Asn Val Ser Gln Thr Asn 85 90 95	288												
ttc aaa agt ctt tta aga aat ctt ggt gtt tca gag aat ata ttc ctt Phe Lys Ser Leu Leu Arg Asn Leu Gly Val Ser Glu Asn Ile Phe Leu 100 105 110	336												
aaa gaa aat aaa tta aat aag gaa ggg aaa tta att gaa cac ata ata Lys Glu Asn Lys Leu Asn Lys Glu Gly Lys Leu Ile Glu His Ile Ile 115 120 125	384												
aat gat gat gac gat aaa aaa tat att aaa ggg caa gac gaa aac Asn Asp Asp Asp Lys Lys Lys Tyr Ile Lys Gly Gln Asp Glu Asn 130 135 140	432												

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			tta Leu								576
_		_	gat Asp	-		_	_	_	_	_	624
			agc Ser								672
			caa Gln 230								720
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- Glu Ile Ile Lys Ser Asn Leu Arg Ser Gly Ser Ser Asn Ser Arg Asn 35 40 45
- Arg Ile Asn Glu Glu Asn His Glu Lys Lys His Val Leu Ser His Asn 50 55 60
- Ser Tyr Glu Lys Thr Lys Asn Asn Glu Asn Asn Lys Phe Phe Asp Lys 65 70 75 80
- Asp Lys Glu Leu Thr Met Ser Asn Val Lys Asn Val Ser Gln Thr Asn 85 90 95
- Phe Lys Ser Leu Leu Arg Asn Leu Gly Val Ser Glu Asn Ile Phe Leu 100 105 110
- Lys Glu Asn Lys Leu Asn Lys Glu Gly Lys Leu Ile Glu His Ile Ile 115 120 125
- Asn Asp Asp Asp Lys Lys Lys Tyr Ile Lys Gly Gln Asp Glu Asn 130 135 140
- Arg Gln Glu Asp Leu Glu Glu Lys Ala Ala Lys Glu Lys Leu Gln Gly 145 150 155 160
- Gln Gln Ser Asp Ser Glu Gln Glu Arg Arg Ala Lys Glu Lys Leu Gln 165 170 175
- Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Leu Ala Lys Glu Lys Leu 180 185 190
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- Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Arg Ala Lys 225 230 235 240
- Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Arg Ala 245 250 255
- Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Leu 260 265 270
- Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Asp Arg 275 280 285
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ctaaagaaaa gttgcaagaa caacaaagcg atttagaaca agatagactt gctaaagaaa 300
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cgatttagaa caagagagac ttgctaaaga aaagttgcaa gaacaacaaa gcgatttaga 480
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				gaa Glu												384
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Leu Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu

Arg Arg Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln 115 120 125

Glu Arg Arg Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu 130 135 140

Gln Glu Arg Leu Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu 145 150 155 160

Glu Gln Glu Arg Arg Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp 165 170 175

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- Arg Asp Leu Glu Gln Arg Lys Ala Asp Thr Lys Lys Asn Leu Glu Arg 210 215 220
- Lys Lys Glu His Gly Asp Ile Leu Ala Glu Asp Leu Tyr Gly Arg Leu 225 230 235 240
- Glu Ile Pro Ala Ile Glu Leu Pro Ser Glu Asn Glu Arg Gly Tyr Tyr
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- Asp Ser Lys Glu Ile Ser Ile Ile Glu Lys Thr Asn Arg Glu Ser Ile 275 280 285
- Thr Thr Asn Val Glu Gly Arg Arg Asp Ile His Lys Gly His Leu Glu 290 295 300
- Glu Lys Lys Asp Gly Ser Ile Lys Pro Glu Gln Lys Glu Asp Lys Ser 305 310 315 320
- Ala Asp Ile Gln Asn His Thr Leu Glu Thr Val Asn Ile Ser Asp Val
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- Gly Ile Tyr Lys Glu Leu Glu Asp Leu Ile Glu Lys Asn Glu Asn Leu 385 390 395 400
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- Glu Lys Ile Lys Lys Gly Lys Lys Tyr Glu Lys Thr Lys Asp Asn Asn 420 425 430
- Phe Lys Pro Asn Asp Lys Ser Leu Tyr Asp Glu His Ile Lys Lys Tyr 435 440 445
- Lys Asn Asp Lys Gln Val Asn Lys Glu Lys Glu Lys Phe Ile Lys Ser 450 455 460
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Leu Gln Glu Gln Ser Asp Leu Glu Gln Asp Arg Leu Ala Lys Glu
aag tta caa gag cag caa agc gat tta gaa caa gag aga ctt gct aaa
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Lys Leu Gln Gln Gln Ser Asp Leu Gln Gln Arg Leu Ala Lys
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Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg Arg
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Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Asp Arg
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Leu Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu
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				caa Gln								1152
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- Leu Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln Glu
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- Arg Arg Ala Lys Glu Lys Leu Gln Glu Gln Gln Ser Asp Leu Glu Gln 115 120 125
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- Phe Lys Pro Ile Val Gln Tyr Asp Asn Phe Gln Asp Glu Glu Asn Ile 370 375 380
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- Lys Asn Asp Lys Gln Val Asn Lys Glu Lys Glu Lys Phe Ile Lys Ser 450 460
- Leu Phe His Ile Phe Asp Gly Asp Asn Glu Ile Leu Gln Ile Val Asp 465 470 470 480
- Glu Leu Ser Glu Asp Ile Thr Lys Tyr Phe Met Lys Leu 485 490